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Substitute for form 1449A/PTO INFORMATION DISCLOSURE STATEMENT BY APPLICANT (use as many sheets as necessary)				Complete if Known	
				Application Number	10/047,724
				Filing Date	January 15, 2002
				First Named Inventor	Vince Hilser
				Art Unit	1631
				Examiner Name	C. D. Ly
Sheet	1	of	1	Attorney Docket Number	HO-P02070US1

U.S. PATENT DOCUMENTS					
Examiner Initials*	Cite No. ¹	Document Number	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear
		Number-Kind Code ² (if known)			
CO	AA	US-20030032065-A1	02-13-2003	Hilser et al.	
	AB	US-10/096,178	10/15/2002	Fox, R. et al.	
	AC	US-20030088392-A1	05-08-2003	Hilser et al.	

FOREIGN PATENT DOCUMENTS						
Examiner Initials*	Cite No. ¹	Foreign Patent Document	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear	T ⁶
		Country Code ³ -Number ⁴ -Kind Code ⁵ (if known)				
CO	BA	WO 98/54665	12/03/1998	The John Hopkins University		
	BB	WO 02/073193 A1	09/19/2002	Board of Regents UT System		

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NON PATENT LITERATURE DOCUMENTS			
Examiner Initials	Cite No. ¹	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc), date, page(s), volume-issue number(s), publisher, city and/or country where published.	T ²
CO	CA	Ferreon, Josephine et al.; The Effect of the polyproline II (PPII) conformation on the denatured state entropy; Protein Science 2003, 12:447-457.	
	CB	Kim, Hwa-OK et al.; A merger of Rational Drug Design and Combinatorial Chemistry: Development and Application of Peptide Secondary Structure Mimetics; Combinational Chemistry & High Throughput Screening, 2000, Vol. 3, No. 3; 168-183.	
	CC	Lengauer, Thomas et al. Protein structure prediction methods for drug design; Briefings in Bioinformatics; 2000; Vol. 1. No. 3. 275-288.	
	CD	Wrabl, James O. et al.; Thermodynamic propensities of amino acids in the native state ensemble: Implications for fold recognition; Protein Science, 2000; 10:1032-1045.	

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